



LUD5826US PTO SEQUENCE LISTING (1 SEQ
SEQUENCE LISTING

<110> GNJATIC, Sacha
ATANACKOVIC, Djordje
OLD, Lloyd J.

<120> Isolated NY-ESO-1 Peptides which Bind to HLA Class II Molecules
and Uses Thereof

<130> LUD 5826 US

<140> 10/556,853
<141> 2008-02-11

<150> PCT/US04/16849
<151> 2004-05-27

<150> 60/474,893
<151> 2003-05-30

<160> 1

<170> PatentIn version 3.2

<210> 1
<211> 180
<212> PRT
<213> Homo sapiens
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Ala	Gly	Gly	Pro	Gly	Glu	Ala	Gly	Ala	Thr	Gly	Gly	Arg	Ala	Pro
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Cys	Arg	Cys	Gly	Ala	Arg	Gly	Pro	Glu	Ser	Arg	Leu	Leu	Glu	Phe
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Tyr	Leu	Ala	Met	Pro	Phe	Ala	Thr	Pro	Met	Glu	Ala	Glu	Leu	Ala
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Arg	Leu	Thr	Ala	Ala	Asp	His	Arg	Gln	Leu	Gln	Leu	Ser	Ile	Ser
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Ser	Cys	Leu	Gln	Gln	Leu	Ser	Leu	Leu	Met	Trp	Ile	Thr	Gln	Cys
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LUD5826US PTO SEQUENCE LISTING (1 SEQ

Phe	Leu	Pro	Val	Phe	Leu	Ala	Gln	Pro	Pro	Ser	Gly	Gln	Arg	Arg
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Sequence Listing could not be accepted due to errors.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).

Reviewer: Keisha Douglas

Timestamp: [year=2011; month=3; day=28; hr=11; min=18; sec=10; ms=496;]

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Reviewer Comments:

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Met Gln Ala Glu Gly Arg Gly Thr Gly Gly Ser Thr Gly Asp Ala

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10

15

Please correct the misaligned amino acid codings in the above
Sequence ID# 1, found at locations 1 through 15.

Application No: 10556853

Version No: 3.0

Input Set:

Output Set:

Started: 2011-03-10 10:31:27.859

Finished: 2011-03-10 10:31:28.300

Elapsed: 0 hr(s) 0 min(s) 0 sec(s) 441 ms

Total Warnings: 0

Total Errors: 4

No. of SeqIDs Defined: 1

Actual SeqID Count: 1

Error code	Error Description
E 323	Invalid/missing amino acid numbering SEQID (1) POS (9)
E 323	Invalid/missing amino acid numbering SEQID (1)at Protein (10)
E 323	Invalid/missing amino acid numbering SEQID (1) POS (14)
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Arg	Gly	Ala	Gly	Ala	Ala	Arg	Ala	Ser	Gly	Pro	Gly	Gly	Gly	Ala	50	55	60	
Pro	Arg	Gly	Pro	His	Gly	Gly	Ala	Ala	Ser	Gly	Leu	Asn	Gly	Cys	65	70	75	
Cys	Arg	Cys	Gly	Ala	Arg	Gly	Pro	Glu	Ser	Arg	Leu	Leu	Glu	Phe	80	85	90	
Tyr	Leu	Ala	Met	Pro	Phe	Ala	Thr	Pro	Met	Glu	Ala	Glu	Leu	Ala	95	100	105	
Arg	Arg	Ser	Leu	Ala	Gln	Asp	Ala	Pro	Pro	Leu	Pro	Val	Pro	Gly	110	115	120	
Val	Leu	Leu	Lys	Glu	Phe	Thr	Val	Ser	Gly	Asn	Ile	Leu	Thr	Ile				

125

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Arg Leu Thr Ala Ala Asp His Arg Gln Leu Gln Leu Ser Ile Ser

140

145

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Ser Cys Leu Gln Gln Leu Ser Leu Leu Met Trp Ile Thr Gln Cys

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Phe Leu Pro Val Phe Leu Ala Gln Pro Pro Ser Gly Gln Arg Arg

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